OIPE

#3

RAW SEQUENCE LISTING　　　　　　　DATE: 12/06/2001
PATENT APPLICATION: US/09/899,634A　　　TIME: 16:19:12 ENTERED

Input Set : A:\SEQ IDs of application 4_31499A.txt
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    3 <110> APPLICANT: Thomas Buehler, Reto Andreas Gadient, Reinhard Korn, Rao Movva
W-->  4 <120> TITLE OF INVENTION: pCAR and its uses
    6 <130> FILE REFERENCE: 4-31499A
C-->  8 <140> CURRENT APPLICATION NUMBER: US/09/899,634A
C-->  8 <141> CURRENT FILING DATE: 2001-07-05
    8 <160> NUMBER OF SEQ ID NOS: 4
   10 <170> SOFTWARE: PatentIn version 3.0
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   14 <212> TYPE: DNA
   15 <213> ORGANISM: porcine
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142                 40                  45                  50
144 ccg ctg gac atc gag tgg ctg ctg tca cca gct gat aat cag aag gtg  3429
145 Pro Leu Asp Ile Glu Trp Leu Leu Ser Pro Ala Asp Asn Gln Lys Val
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149 Asp Gln Val Ile Ile Leu Tyr Ser Gly Asp Lys Ile Tyr Asp Asp Tyr
150         70                  75                  80
152 tac caa gat ctg aaa gga cga gta cat ttt aca agt aat gat ctc aaa  3525
153 Tyr Gln Asp Leu Lys Gly Arg Val His Phe Thr Ser Asn Asp Leu Lys
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157 Ser Gly Asp Ala Ser Ile Asn Val Thr Asn Leu Gln Leu Ser Asp Ile
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173 Pro Lys Glu Gly Ser Leu Pro Leu Leu Tyr Glu Trp Gln Lys Leu Ser
174     165             170             175
176 aat tca cag aag ctg ccc acc ttg tgg tta gca gaa atg act tca cct      3813
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178 180             185             190             195
180 gtt ata tct gta aaa aat gcc tct act gaa tac tct ggg aca tac agc      3861
181 Val Ile Ser Val Lys Asn Ala Ser Thr Glu Tyr Ser Gly Thr Tyr Ser
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184 tgt acc gtg aaa aac aga gtg ggc tct gat cag tgc ctg ctt cgc ctg      3909
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193 Ile Gly Val Leu Leu Ala Leu Val Leu Ile Gly Leu Ile Ile Phe Cys
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197 Cys Arg
198 260
200 tagaggtttt acttgcttta aaaaacctcc cacacctccc cctgaacctg aaacataaaa   4114
202 tgaatgcaat tgttgttgtt aacttgttta ttgcagctta taatggttac aaataaaagca  4174
204 atagcatcac aaatttcaca aataaagcat ttttttcact gcattctagt tgtggtttgt   4234
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224 Lys Gly Glu Thr Ala Tyr Leu Pro Cys Arg Phe Thr Leu Gly Pro Glu
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228 Asp Gln Gly Pro Leu Asp Ile Glu Trp Leu Leu Ser Pro Ala Asp Asn
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232 Gln Lys Val Asp Gln Val Ile Ile Leu Tyr Ser Gly Asp Lys Ile Tyr
233 65             70             75             80
236 Asp Asp Tyr Tyr Gln Asp Leu Lys Gly Arg Val His Phe Thr Ser Asn
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252 Arg Cys Tyr Val Asp Gly Ser Glu Glu Ile Gly Asn Asp Phe Lys Leu
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256 Lys Cys Glu Pro Lys Glu Gly Ser Leu Pro Leu Leu Tyr Glu Trp Gln
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261             180                 185                 190
264 Thr Ser Pro Val Ile Ser Val Lys Asn Ala Ser Thr Glu Tyr Ser Gly
265         195                 200                 205
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269     210                 215                 220
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303 aaa ggg gaa act gcc tat ttg cca tgc aga ttt acc ctg ggt cca gaa       144
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361                 260                 265                 270
363 gtg cat cat gat atc agg gaa gac gtg cct cct ccg aag agc aga acg     864
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377                 325                 330                 335
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